SEQUENCE LISTING

<110> Williams, Kevin J.

<120> Thrombospondin Fragments and Uses Thereof In Clinical Assays for Cancer and Generation of Antibodies and Other Binding Agents

<130> W1107-20005

<140> 10/419,462
<141> 2003-04-21

<160> 64

<170> PatentIn version 3.3

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<400> 1

Thr Glu Glu Asn Lys Glu
1 5

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<220>
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<400> 2

Cys Leu Gln Asp Ser Ile Arg Lys Val Thr Glu Glu Asn Lys Glu
1 5 10 15

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<400> 3

Leu Gln Asp Ser Ile Arg Lys Val Thr Glu Glu Asn Lys Glu
1 5 10

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Glu Gly Glu Ala Arg Glu
1 5

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<400> 5

Pro Gln Met Asn Gly Lys Pro Cys Glu Gly Glu Ala Arg Glu
1 5 10

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<400> 6

Glu Asp Thr Asp Leu Asp
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<400> 7

Tyr Ala Gly Asn Gly Ile Ile Cys Gly Glu Asp Thr Asp Leu Asp
1 5 10 15

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<220>
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<400> 8

Cys Asn Ser Pro Ser Pro Gln Met Asn Gly Lys Pro Cys Glu Gly Glu

1 5 10 15

Ala Arg

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<220>
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<400> 9

Arg Lys Val Thr Glu Glu Asn Lys Glu Leu Ala Asn Glu Leu Arg Arg
1 5 10 15

Pro

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<220>
<223> Thrombospondin Region which includes an N-terminal CYS added to aid conjugation

<400> 10

Cys Arg Lys Val Thr Glu Glu Asn Lys Glu Leu Ala Asn Glu Leu Arg
1 5 10 15

Arg Pro

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<400> 11

Pro Gln Met Asn Gly Lys Pro Cys Glu Gly Glu Ala Arg
1 5 10

<210> 12
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<400> 12

Cys Glu Gly Glu Ala Arg
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<211> 9
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<400> 13

Arg Lys Val Thr Glu Glu Asn Lys Glu
1 5

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<400> 14

Asp Asp Asp Asp Asn Asp Lys Ile Pro Asp Asp Arg Asp Asn Cys
1 5 10 15

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<400> 15

Asp Asp Asp Asp Asn Asp Lys Ile Pro Asp Asp Arg Asp Asn Cys
1 5 10 15

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Asp Asp Asp Asp Asn Asp Lys
1 5

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<400> 17

Asn Leu Pro Asn Ser Gly Gln Glu Asp Tyr Asp Lys Asp Gly
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<400> 18

Cys Asn Leu Pro Asn Ser Gly Gln Glu Asp Tyr Asp Lys Asp Gly
1 5 10 15

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<400> 19

Glu Asp Tyr Asp Lys Asp
1 5

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<400> 20

Cys Pro Tyr Asn His Asn Pro Asp Gln Ala Asp Thr Asp Asn Asn Gly
1 5 10 15

Glu Gly Asp

<210> 21

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<400> 21

Cys Arg Leu Val Pro Asn Pro Asp Gln Lys Asp Ser Asp Gly Asp
1 5 10 15

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<400> 22

Asp Gln Lys Asp Ser Asp Gly Asp
1 5

<210> 23
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<400> 23

Cys Pro Tyr Val Pro Asn Ala Asn Gln Ala Asp His Asp Lys Asp Gly
1 5 10 15

Lys Gly Asp Ala
20

<210> 24
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<400> 24

Thr Glu Arg Asp Asp Asp
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<210> 25
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<400> 25

Asp Phe Ser Gly Thr Phe Phe Ile Asn Thr Glu Arg Asp Asp Asp
1 5 10 15

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<400> 26

Glu Arg Lys Asp His Ser
1 5

<210> 27
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<220>
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<400> 27

Thr Arg Gly Thr Leu Leu Ala Leu Glu Arg Lys Asp His Ser
1 5 10

<210> 28
<211> 15
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<223> Thrombospondin Region plus N-termianl CYS

<400> 28

Cys Thr Arg Gly Thr Leu Leu Ala Leu Glu Arg Lys Asp His Ser
1 5 10 15

<210> 29
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<400> 29

Asp Asp Lys Phe Gln Asp

1 5

<210> 30
<211> 14
<212> PRT
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<220>
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<400> 30

Ala Asn Leu Ile Pro Pro Val Pro Asp Asp Lys Phe Gln Asp
1 5 10

<210> 31
<211> 15
<212> PRT
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<220>
<223> Thrombospondin Region plus N-terminal CYS

<400> 31

Cys Ala Asn Leu Ile Pro Pro Val Pro Asp Asp Lys Phe Gln Asp
1 5 10 15

<210> 32
<211> 6
<212> PRT
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<220>
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<400> 32

Asp Cys Glu Lys Met Glu
1 5

<210> 33
<211> 15
<212> PRT
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<223> Thrombospondin Region

<400> 33

Glu Asp Arg Ala Gln Leu Tyr Ile Asp Cys Glu Lys Met Glu Asn
1 5 10 15

<210> 34
<211> 17
<212> PRT
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<220>
<223> Thrombospondin Region

<400> 34

Cys Gly Thr Asn Arg Ile Pro Glu Ser Gly Gly Asp Asn Ser Val Phe
1 5 10 15

Asp

<210> 35
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<220>
<223> Thrombospondin region

<400> 35

Asn Arg Ile Pro Glu Ser Gly Gly Asp Asn Ser Val Phe Asp
1 5 10

<210> 36
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<212> PRT
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<220>
<223> Thrombospondin Region

<400> 36

Gly Trp Lys Asp Phe Thr Ala Tyr Arg Trp Arg Leu Ser His Arg Pro
1 5 10 15

Lys Thr Gly

<210> 37
<211> 20
<212> PRT
<213> Artificial

<220>
<223> Thrombospondin region plus N-terminal CYS

<400> 37

Cys Gly Trp Lys Asp Phe Thr Ala Tyr Arg Trp Arg Leu Ser His Arg
1 5 10 15

Pro Lys Thr Gly
20

<210> 38
<211> 1170
<212> PRT
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<400> 38

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Met Gly Leu Ala Trp Gly Leu Gly Val Leu Phe Leu Met His Val Cys
1               5                  10                  15

Gly Thr Asn Arg Ile Pro Glu Ser Gly Gly Asp Asn Ser Val Phe Asp
            20                  25                  30

Ile Phe Glu Leu Thr Gly Ala Ala Arg Lys Gly Ser Gly Arg Arg Leu
        35                  40                  45

Val Lys Gly Pro Asp Pro Ser Ser Pro Ala Phe Arg Ile Glu Asp Ala
    50                  55                  60

Asn Leu Ile Pro Pro Val Pro Asp Asp Lys Phe Gln Asp Leu Val Asp
65                  70                  75                  80

Ala Val Arg Ala Glu Lys Gly Phe Leu Leu Leu Ala Ser Leu Arg Gln
                85                  90                  95

Met Lys Lys Thr Arg Gly Thr Leu Leu Ala Leu Glu Arg Lys Asp His
            100                 105                 110

Ser Gly Gln Val Phe Ser Val Val Ser Asn Gly Lys Ala Gly Thr Leu
        115                 120                 125

Asp Leu Ser Leu Thr Val Gln Gly Lys Gln His Val Val Ser Val Glu
    130                 135                 140

Glu Ala Leu Leu Ala Thr Gly Gln Trp Lys Ser Ile Thr Leu Phe Val
145                 150                 155                 160

Gln Glu Asp Arg Ala Gln Leu Tyr Ile Asp Cys Glu Lys Met Glu Asn
                165                 170                 175

Ala Glu Leu Asp Val Pro Ile Gln Ser Val Phe Thr Arg Asp Leu Ala
            180                 185                 190

Ser Ile Ala Arg Leu Arg Ile Ala Lys Gly Gly Val Asn Asp Asn Phe
        195                 200                 205

Gln Gly Val Leu Gln Asn Val Arg Phe Val Phe Gly Thr Thr Pro Glu
    210                 215                 220
```

```
Asp Ile Leu Arg Asn Lys Gly Cys Ser Ser Ser Thr Ser Val Leu Leu
225                 230                 235                 240

Thr Leu Asp Asn Asn Val Val Asn Gly Ser Ser Pro Ala Ile Arg Thr
                245                 250                 255

Asn Tyr Ile Gly His Lys Thr Lys Asp Leu Gln Ala Ile Cys Gly Ile
            260                 265                 270

Ser Cys Asp Glu Leu Ser Ser Met Val Leu Glu Leu Arg Gly Leu Arg
        275                 280                 285

Thr Ile Val Thr Thr Leu Gln Asp Ser Ile Arg Lys Val Thr Glu Glu
    290                 295                 300

Asn Lys Glu Leu Ala Asn Glu Leu Arg Arg Pro Pro Leu Cys Tyr His
305                 310                 315                 320

Asn Gly Val Gln Tyr Arg Asn Asn Glu Glu Trp Thr Val Asp Ser Cys
                325                 330                 335

Thr Glu Cys His Cys Gln Asn Ser Val Thr Ile Cys Lys Lys Val Ser
            340                 345                 350

Cys Pro Ile Met Pro Cys Ser Asn Ala Thr Val Pro Asp Gly Glu Cys
        355                 360                 365

Cys Pro Arg Cys Trp Pro Ser Asp Ser Ala Asp Asp Gly Trp Ser Pro
    370                 375                 380

Trp Ser Glu Trp Thr Ser Cys Ser Thr Ser Cys Gly Asn Gly Ile Gln
385                 390                 395                 400

Gln Arg Gly Arg Ser Cys Asp Ser Leu Asn Asn Arg Cys Glu Gly Ser
                405                 410                 415

Ser Val Gln Thr Arg Thr Cys His Ile Gln Glu Cys Asp Lys Arg Phe
            420                 425                 430

Lys Gln Asp Gly Gly Trp Ser His Trp Ser Pro Trp Ser Ser Cys Ser
        435                 440                 445

Val Thr Cys Gly Asp Gly Val Ile Thr Arg Ile Arg Leu Cys Asn Ser
    450                 455                 460

Pro Ser Pro Gln Met Asn Gly Lys Pro Cys Glu Gly Glu Ala Arg Glu
465                 470                 475                 480
```

```
Thr Lys Ala Cys Lys Lys Asp Ala Cys Pro Ile Asn Gly Gly Trp Gly
                485                 490                 495

Pro Trp Ser Pro Trp Asp Ile Cys Ser Val Thr Cys Gly Gly Gly Val
            500                 505                 510

Gln Lys Arg Ser Arg Leu Cys Asn Asn Pro Ala Pro Gln Phe Gly Gly
        515                 520                 525

Lys Asp Cys Val Gly Asp Val Thr Glu Asn Gln Ile Cys Asn Lys Gln
    530                 535                 540

Asp Cys Pro Ile Asp Gly Cys Leu Ser Asn Pro Cys Phe Ala Gly Val
545                 550                 555                 560

Lys Cys Thr Ser Tyr Pro Asp Gly Ser Trp Lys Cys Gly Ala Cys Pro
                565                 570                 575

Pro Gly Tyr Ser Gly Asn Gly Ile Gln Cys Thr Asp Val Asp Glu Cys
            580                 585                 590

Lys Glu Val Pro Asp Ala Cys Phe Asn His Asn Gly Glu His Arg Cys
        595                 600                 605

Glu Asn Thr Asp Pro Gly Tyr Asn Cys Leu Pro Cys Pro Pro Arg Phe
    610                 615                 620

Thr Gly Ser Gln Pro Phe Gly Gln Gly Val Glu His Ala Thr Ala Asn
625                 630                 635                 640

Lys Gln Val Cys Lys Pro Arg Asn Pro Cys Thr Asp Gly Thr His Asp
                645                 650                 655

Cys Asn Lys Asn Ala Lys Cys Asn Tyr Leu Gly His Tyr Ser Asp Pro
            660                 665                 670

Met Tyr Arg Cys Glu Cys Lys Pro Gly Tyr Ala Gly Asn Gly Ile Ile
        675                 680                 685

Cys Gly Glu Asp Thr Asp Leu Asp Gly Trp Pro Asn Glu Asn Leu Val
    690                 695                 700

Cys Val Ala Asn Ala Thr Tyr His Cys Lys Lys Asp Asn Cys Pro Asn
705                 710                 715                 720

Leu Pro Asn Ser Gly Gln Glu Asp Tyr Asp Lys Asp Gly Ile Gly Asp
```

725 730 735

Ala Cys Asp Asp Asp Asp Asp Asn Asp Lys Ile Pro Asp Asp Arg Asp
740 745 750

Asn Cys Pro Phe His Tyr Asn Pro Ala Gln Tyr Asp Tyr Asp Arg Asp
755 760 765

Asp Val Gly Asp Arg Cys Asp Asn Cys Pro Tyr Asn His Asn Pro Asp
770 775 780

Gln Ala Asp Thr Asp Asn Asn Gly Glu Gly Asp Ala Cys Ala Ala Asp
785 790 795 800

Ile Asp Gly Asp Gly Ile Leu Asn Glu Arg Asp Asn Cys Gln Tyr Val
805 810 815

Tyr Asn Val Asp Gln Arg Asp Thr Asp Met Asp Gly Val Gly Asp Gln
820 825 830

Cys Asp Asn Cys Pro Leu Glu His Asn Pro Asp Gln Leu Asp Ser Asp
835 840 845

Ser Asp Arg Ile Gly Asp Thr Cys Asp Asn Asn Gln Asp Ile Asp Glu
850 855 860

Asp Gly His Gln Asn Asn Leu Asp Asn Cys Pro Tyr Val Pro Asn Ala
865 870 875 880

Asn Gln Ala Asp His Asp Lys Asp Gly Lys Gly Asp Ala Cys Asp His
885 890 895

Asp Asp Asp Asn Asp Gly Ile Pro Asp Asp Lys Asp Asn Cys Arg Leu
900 905 910

Val Pro Asn Pro Asp Gln Lys Asp Ser Asp Gly Asp Gly Arg Gly Asp
915 920 925

Ala Cys Lys Asp Asp Phe Asp His Asp Ser Val Pro Asp Ile Asp Asp
930 935 940

Ile Cys Pro Glu Asn Val Asp Ile Ser Glu Thr Asp Phe Arg Arg Phe
945 950 955 960

Gln Met Ile Pro Leu Asp Pro Lys Gly Thr Ser Gln Asn Asp Pro Asn
965 970 975

```
Trp Val Val Arg His Gln Gly Lys Glu Leu Val Gln Thr Val Asn Cys
            980                 985                 990

Asp Pro Gly Leu Ala Val Gly Tyr  Asp Glu Phe Asn Ala  Val Asp Phe
        995                 1000                1005

Ser Gly  Thr Phe Phe Ile Asn  Thr Glu Arg Asp Asp  Asp Tyr Ala
    1010                1015                1020

Gly Phe  Val Phe Gly Tyr Gln  Ser Ser Ser Arg Phe  Tyr Val Val
    1025                1030                1035

Met Trp  Lys Gln Val Thr Gln  Ser Tyr Trp Asp Thr  Asn Pro Thr
    1040                1045                1050

Arg Ala  Gln Gly Tyr Ser Gly  Leu Ser Val Lys Val  Val Asn Ser
    1055                1060                1065

Thr Thr  Gly Pro Gly Glu His  Leu Arg Asn Ala Leu  Trp His Thr
    1070                1075                1080

Gly Asn  Thr Pro Gly Gln Val  Arg Thr Leu Trp His  Asp Pro Arg
    1085                1090                1095

His Ile  Gly Trp Lys Asp Phe  Thr Ala Tyr Arg Trp  Arg Leu Ser
    1100                1105                1110

His Arg  Pro Lys Thr Gly Phe  Ile Arg Val Val Met  Tyr Glu Gly
    1115                1120                1125

Lys Lys  Ile Met Ala Asp Ser  Gly Pro Ile Tyr Asp  Lys Thr Tyr
    1130                1135                1140

Ala Gly  Gly Arg Leu Gly Leu  Phe Val Phe Ser Gln  Glu Met Val
    1145                1150                1155

Phe Phe  Ser Asp Leu Lys Tyr  Glu Cys Arg Asp Pro
    1160                1165                1170


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<400>  39

Met Gly Leu Ala Trp Gly Leu Gly Val Leu Phe Leu Met His Val Cys
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1 5 10 15

Gly Thr

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<400> 40

Asn Arg Ile Pro Glu Ser Gly Gly Asp Asn Ser Val Phe Asp Ile Phe
1 5 10 15

Glu Leu Thr Gly Ala Ala Arg Lys Gly Ser Gly Arg Arg Leu Val Lys
20 25 30

Gly Pro Asp Pro Ser Ser Pro Ala Phe Arg Ile Glu Asp Ala Asn Leu
35 40 45

Ile Pro Pro Val Pro Asp Asp Lys Phe Gln Asp Leu Val Asp Ala Val
50 55 60

Arg Ala Glu Lys Gly Phe Leu Leu Leu Ala Ser Leu Arg Gln Met Lys
65 70 75 80

Lys Thr Arg Gly Thr Leu Leu Ala Leu Glu Arg Lys Asp His Ser Gly
85 90 95

Gln Val Phe Ser Val Val Ser Asn Gly Lys Ala Gly Thr Leu Asp Leu
100 105 110

Ser Leu Thr Val Gln Gly Lys Gln His Val Val Ser Val Glu Glu Ala
115 120 125

Leu Leu Ala Thr Gly Gln Trp Lys Ser Ile Thr Leu Phe Val Gln Glu
130 135 140

Asp Arg Ala Gln Leu Tyr Ile Asp Cys Glu Lys Met Glu Asn Ala Glu
145 150 155 160

Leu Asp Val Pro Ile Gln Ser Val Phe Thr Arg Asp Leu Ala Ser Ile
165 170 175

Ala Arg Leu Arg Ile Ala Lys Gly Gly Val Asn Asp Asn Phe Gln Gly
180 185 190

Val Leu Gln Asn Val Arg Phe Val Phe Gly Thr Thr Pro Glu Asp Ile
195 200 205

Leu Arg Asn Lys Gly Cys Ser Ser Ser Thr Ser Val Leu Leu Thr Leu
210 215 220

Asp Asn Asn Val Val Asn Gly Ser Ser Pro Ala Ile Arg Thr Asn Tyr
225 230 235 240

<210> 41
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<223> Thrombospondin region plus domain of inter-chain disulfide bonds

<400> 41

Ile Gly His Lys Thr Lys Asp Leu Gln Ala Ile Cys Gly Ile Ser Cys
1 5 10 15

Asp Glu Leu Ser Ser Met
20

<210> 42
<211> 98
<212> PRT
<213> Artificial

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<223> Thrombospondin region plus procollagen homology domain

<400> 42

Val Leu Glu Leu Arg Gly Leu Arg Thr Ile Val Thr Thr Leu Gln Asp
1 5 10 15

Ser Ile Arg Lys Val Thr Glu Glu Asn Lys Glu Leu Ala Asn Glu Leu
20 25 30

Arg Arg Pro Pro Leu Cys Tyr His Asn Gly Val Gln Tyr Arg Asn Asn
35 40 45

Glu Glu Trp Thr Val Asp Ser Cys Thr Glu Cys His Cys Gln Asn Ser
50 55 60

Val Thr Ile Cys Lys Lys Val Ser Cys Pro Ile Met Pro Cys Ser Asn
65 70 75 80

Ala Thr Val Pro Asp Gly Glu Cys Cys Pro Arg Cys Trp Pro Ser Asp

85 90 95

Ser Ala

<210> 43
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<223> Thrombospondin region plus domain of type 1 repeats

<400> 43

Asp Asp Gly Trp Ser Pro Trp Ser Glu Trp Thr Ser Cys Ser Thr Ser
1 5 10 15

Cys Gly Asn Gly Ile Gln Gln Arg Gly Arg Ser Cys Asp Ser Leu Asn
20 25 30

Asn Arg Cys Glu Gly Ser Ser Val Gln Thr Arg Thr Cys His Ile Gln
35 40 45

Glu Cys Asp Lys Arg Phe Lys Gln Asp Gly Gly Trp Ser His Trp Ser
50 55 60

Pro Trp Ser Ser Cys Ser Val Thr Cys Gly Asp Gly Val Ile Thr Arg
65 70 75 80

Ile Arg Leu Cys Asn Ser Pro Ser Pro Gln Met Asn Gly Lys Pro Cys
85 90 95

Glu Gly Glu Ala Arg Glu Thr Lys Ala Cys Lys Lys Asp Ala Cys Pro
100 105 110

Ile Asn Gly Gly Trp Gly Pro Trp Ser Pro Trp Asp Ile Cys Ser Val
115 120 125

Thr Cys Gly Gly Gly Val Gln Lys Arg Ser Arg Leu Cys Asn Asn Pro
130 135 140

Ala Pro Gln Phe Gly Gly Lys Asp Cys Val Gly Asp Val Thr Glu Asn
145 150 155 160

Gln Ile Cys Asn Lys Gln Asp Cys Pro Ile
165 170

<210> 44
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<212> PRT
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<223> Thrombospondin region plus domain of type 2 repeats

<400> 44

Asp Gly Cys Leu Ser Asn Pro Cys Phe Ala Gly Val Lys Cys Thr Ser
1 5 10 15

Tyr Pro Asp Gly Ser Trp Lys Cys Gly Ala Cys Pro Pro Gly Tyr Ser
20 25 30

Gly Asn Gly Ile Gln Cys Thr Asp Val Asp Glu Cys Lys Glu Val Pro
35 40 45

Asp Ala Cys Phe Asn His Asn Gly Glu His Arg Cys Glu Asn Thr Asp
50 55 60

Pro Gly Tyr Asn Cys Leu Pro Cys Pro Pro Arg Phe Thr Gly Ser Gln
65 70 75 80

Pro Phe Gly Gln Gly Val Glu His Ala Thr Ala Asn Lys Gln Val Cys
85 90 95

Lys Pro Arg Asn Pro Cys Thr Asp Gly Thr His Asp Cys Asn Lys Asn
100 105 110

Ala Lys Cys Asn Tyr Leu Gly His Tyr Ser Asp Pro Met Tyr Arg Cys
115 120 125

Glu Cys Lys Pro Gly Tyr Ala Gly Asn Gly Ile Ile Cys Gly Glu
130 135 140

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<220>
<223> Thrombospondin Region plus region between the type 2 and the type 3 repeat

<400> 45

Asp Thr Asp Leu Asp Gly Trp Pro Asn Glu Asn Leu Val Cys Val Ala
1 5 10 15

Asn Ala Thr Tyr His Cys Lys Lys
20

<210> 46
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Thrombospondin region

<400> 46

Asp Asn Cys Pro Asn Leu Pro Asn Ser Gly Gln Glu Asp Tyr Asp Lys
1 5 10 15

Asp Gly Ile Gly Asp Ala Cys Asp Asp Asp Asp Asp Asn Asp Lys Ile
20 25 30

Pro Asp Asp Arg
35

<210> 47
<211> 23
<212> PRT
<213> Artificial

<220>
<223> Thrombospondin Region

<400> 47

Asp Asn Cys Pro Phe His Tyr Asn Pro Ala Gln Tyr Asp Tyr Asp Arg
1 5 10 15

Asp Asp Val Gly Asp Arg Cys
20

<210> 48
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Thrombospondin Region

<400> 48

Asp Asn Cys Pro Tyr Asn His Asn Pro Asp Gln Ala Asp Thr Asp Asn
1 5 10 15

Asn Gly Glu Gly Asp Ala Cys Ala Ala Asp Ile Asp Gly Asp Gly Ile
20 25 30

Leu Asn Glu Arg
35

<210> 49

<211> 23
<212> PRT
<213> Artificial

<220>
<223> thrombospondin region

<400> 49

Asp Asn Cys Gln Tyr Val Tyr Asn Val Asp Gln Arg Asp Thr Asp Met
1 5 10 15

Asp Gly Val Gly Asp Gln Cys
20

<210> 50
<211> 38
<212> PRT
<213> Artificial

<220>
<223> Thrombospondin Region

<400> 50

Asp Asn Cys Pro Leu Glu His Asn Pro Asp Gln Leu Asp Ser Asp Ser
1 5 10 15

Asp Arg Ile Gly Asp Thr Cys Asp Asn Asn Gln Asp Ile Asp Glu Asp
20 25 30

Gly His Gln Asn Asn Leu
35

<210> 51
<211> 36
<212> PRT
<213> Artificial

<220>
<223> Thrombospondin region

<400> 51

Asp Asn Cys Pro Tyr Val Pro Asn Ala Asn Gln Ala Asp His Asp Lys
1 5 10 15

Asp Gly Lys Gly Asp Ala Cys Asp His Asp Asp Asp Asn Asp Gly Ile
20 25 30

Pro Asp Asp Lys
35

<210> 52
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<223> Thrombospondin region plus domain of type 3 repeats

<400> 52

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Asp Asn Cys Arg Leu Val Pro Asn Pro Asp Gln Lys Asp Ser Asp Gly
1               5                   10                  15

Asp Gly Arg Gly Asp Ala Cys Lys Asp Asp Phe Asp His Asp Ser Val
            20                  25                  30

Pro Asp Ile Asp
        35
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<210> 53
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<220>
<223> Thrombospondin Region plus C-terminal domain

<400> 53

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Asp Ile Cys Pro Glu Asn Val Asp Ile Ser Glu Thr Asp Phe Arg Arg
1               5                   10                  15

Phe Gln Met Ile Pro Leu Asp Pro Lys Gly Thr Ser Gln Asn Asp Pro
            20                  25                  30

Asn Trp Val Val Arg His Gln Gly Lys Glu Leu Val Gln Thr Val Asn
        35                  40                  45

Cys Asp Pro Gly Leu Ala Val Gly Tyr Asp Glu Phe Asn Ala Val Asp
    50                  55                  60

Phe Ser Gly Thr Phe Phe Ile Asn Thr Glu Arg Asp Asp Asp Tyr Ala
65                  70                  75                  80

Gly Phe Val Phe Gly Tyr Gln Ser Ser Ser Arg Phe Tyr Val Val Met
                85                  90                  95

Trp Lys Gln Val Thr Gln Ser Tyr Trp Asp Thr Asn Pro Thr Arg Ala
            100                 105                 110

Gln Gly Tyr Ser Gly Leu Ser Val Lys Val Val Asn Ser Thr Thr Gly
        115                 120                 125

Pro Gly Glu His Leu Arg Asn Ala Leu Trp His Thr Gly Asn Thr Pro
```

130 135 140

Gly Gln Val Arg Thr Leu Trp His Asp Pro Arg His Ile Gly Trp Lys
145 150 155 160

Asp Phe Thr Ala Tyr Arg Trp Arg Leu Ser His Arg Pro Lys Thr Gly
165 170 175

Phe Ile Arg Val Val Met Tyr Glu Gly Lys Lys Ile Met Ala Asp Ser
180 185 190

Gly Pro Ile Tyr Asp Lys Thr Tyr Ala Gly Gly Arg Leu Gly Leu Phe
195 200 205

Val Phe Ser Gln Glu Met Val Phe Phe Ser Asp Leu Lys Tyr Glu Cys
210 215 220

Arg Asp Pro
225

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<400> 54

Cys Ser Val Thr Cys Gly
1 5

<210> 55
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<400> 55

Arg Phe Tyr Val Val Met Trp Lys
1 5

<210> 56
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Arg Leu Arg Ile Ala Lys Gly Gly Val Asn Asp Asn
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